

Figure 1 Gregg, R.K., et. al.

91A₃H

_____ D _____

SEQ ID NO. 5	TYR	PHE	CYS	ALA	ARG	SER	TYR	SER	GLY	ASP	MET	TYR	CYS	SEQ ID NO. 6	PHE	ASP	TYR	TRP
SEQ ID NO. 7	TAT	TTC	TGT	GCA	AGA	TCG	TAT	TAC	TCT	GGT	GAT	ATG	TAC	SEQ ID NO. 8	TTT	GAC	TAC	TGG

91A₃H-INSβ

_____ INSβ _____

SEQ ID NO. 9	-	-	-	-	-	-	HIS	LEU	VAL	GLU	ALA	LEU	-	LEU	VAL	CYS	GLY	GLU	ARG	GLY	-	-	-	
SEQ ID NO. 10	-	-	-	-	-	-	AGC	C-C	CTA	GTG	-AG	-CG	C-T	---	CT-	GTT	TGC	GGT	GAA	AGA	GGT	---	---	---

91A₃H-HEL

_____ HEL _____

SEQ ID NO. 11	-	-	-	-	-	-	ALA	MET	LYS	ARG	HIS	GLY	LEU	ASP	ASN	TYR	ARG	GLY	TYR	SER	LEU	-	-	-
SEQ ID NO. 12	-	-	-	-	-	-	GCA	ATG	A-G	CGC	CAC	-GG	A-A	G--	AA-	TAT	CGG	GGA	TAT	AGC	CTC	---	---	---

